SESSION RESUMED IN FILE 'AGRICOLA, BIOSIS, CAPLUS, CABA' AT 10:19:23 ON 18 OCT 2001

09/537,654

=> s rad51

L4 1203 RAD51

=> s 14 and maize

L5 8 L4 AND MAIZE

=> d ti 1-8

- L5 ANSWER 1 OF 8 AGRICOLA
- ${\tt TI}$  Three-dimensional microscopy of the  ${\tt Rad51}$  recombination protein during meiotic prophase.
- L5 ANSWER 2 OF 8 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Three-dimensional microscopy of the Rad51 recombination protein during meiotic prophase.
- L5 ANSWER 3 OF 8 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Rad51 distribution is altered in desynaptic2, a maize meiotic mutant that has abnormal chromosome pairing.
- L5 ANSWER 4 OF 8 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Using 3-D immunocytology to explore the roles of mismatch binding proteins in meiosis.
- L5 ANSWER 5 OF 8 CAPLUS COPYRIGHT 2001 ACS
- TI Protein and cDNA sequences of maize RAD51-like gene and uses thereof in transgenic plants
- L5 ANSWER 6 OF 8 CAPLUS COPYRIGHT 2001 ACS
- TI Nucleotide sequences encoding maize RAD51 recombinases
- L5 ANSWER 7 OF 8 CAPLUS COPYRIGHT 2001 ACS
- $\mbox{TI}$   $\,$  Three-dimensional microscopy of the  $\bf Rad51$  recombination protein during meiotic prophase
- L5 ANSWER 8 OF 8 CABA COPYRIGHT 2001 CABI
- $\mbox{TI}$   $\,$  Three-dimensional microscopy of the Rad51 recombination protein during meiotic prophase.

=> d bib abs 6

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L5 ANSWER 6 OF 8 CAPLUS COPYRIGHT 2001 ACS
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AN 1999:529275 CAPLUS

DN 131:154494

TI Nucleotide sequences encoding maize RAD51 recombinases

- IN Bowen, Benjamin A.; Chamberlin, Mark A.; Drummond, Bruce J.; Mcelver, John A.; Rothstein, Rodney J.
- PA Pioneer Hi-Bred International, Inc., USA

SO PCT Int. Appl., 66 pp.

CODEN: PIXXD2

DT Patent

LA English

| FAN.   | CNT             | 1          |     |             |             |      |                 |                |               |                |      |     |          |     |          |     |     |     |
|--------|-----------------|------------|-----|-------------|-------------|------|-----------------|----------------|---------------|----------------|------|-----|----------|-----|----------|-----|-----|-----|
|        | PATENT NO.      |            |     | KIND DATE   |             |      | APPLICATION NO. |                |               |                | DATE |     |          |     |          |     |     |     |
|        |                 |            |     |             |             |      |                 |                |               | _              |      |     |          |     |          |     |     |     |
| ΡI     | WO 9941394      |            |     | Al 19990819 |             | 0819 |                 | WO 1999-US2900 |               |                |      | 0   | 19990211 |     |          |     |     |     |
|        |                 | W:         | ΑL, | ΑM,         | ΑT,         | ΑU,  | AZ,             | BA,            | BB,           | BG,            | BR,  | BY, | CA,      | CH, | CN,      | CU, | CZ, | DE, |
|        |                 |            | DK, | EE,         | ES,         | FΙ,  | GB,             | GE,            | GH,           | GM,            | HR,  | HU, | ID,      | IL, | IN,      | IS, | JP, | KE, |
|        |                 |            | KG, | ΚP,         | ΚR,         | ΚZ,  | LC,             | LK,            | LR,           | LS,            | LT,  | LU, | LV,      | MD, | MG,      | MK, | MN, | MW, |
|        |                 |            | MX, | NO,         | ΝZ,         | PL,  | PT,             | RO,            | RU,           | SD,            | SE,  | SG, | SI,      | SK, | SL,      | TJ, | TM, | TR, |
|        |                 |            | TT, | UA,         | UG,         | UZ,  | VN,             | YU,            | ZW,           | AM,            | ΑZ,  | BY, | KG,      | ΚZ, | MD,      | RU, | TJ, | TM  |
|        |                 | RW:        | GH, | GM,         | ΚE,         | LS,  | MW,             | SD,            | SZ,           | UG,            | ZW,  | AT, | BE,      | CH, | CY,      | DE, | DK, | ES, |
|        |                 |            |     |             |             |      |                 |                |               |                |      |     |          |     | ВJ,      |     |     |     |
|        |                 |            |     | GΑ,         | GN,         | GW,  | ML,             | MR,            | NE,           | SN,            | TD,  | TG  |          |     |          |     |     |     |
|        | AU 9926699      |            |     | A1 19990830 |             |      |                 |                | AU 1999-26699 |                |      |     | 19990211 |     |          |     |     |     |
|        | EΡ              | EP 1053339 |     |             | Al 20001122 |      |                 |                |               | EP 1999-906894 |      |     |          | 4   | 19990211 |     |     |     |
|        |                 | R:         | AT, | BE,         | CH,         | DE,  | DK,             | ES,            | FR,           | GB,            | GR,  | IT, | LI,      | LU, | NL,      | SE, | MC, | PT, |
| IE, FI |                 |            |     |             |             |      |                 |                | •             |                |      |     |          |     |          |     |     |     |
| PRAI   | I US 1998-74745 |            |     | P 19980213  |             |      |                 |                |               |                |      |     |          |     |          |     |     |     |

WO 1999-US2900 W 19990211

AB Nucleic acid sequences encoding two RAD51 recombinases active in maize plants are provided. Full-length cDNA sequences from a maize tassel cDNA library, including the ZmRAD51 coding sequences and unique 3-untranslated regions which are useful as RFLP probes, are

also provided. RFLP mapping indicates that the Zea mays genome contains 2 genes encoding different variants of the ZmRAD51 recombinase enzyme, one mapped to chromosome 7 and the other on chromosome 3. The prodn. of plasmids contg. a nucleic acid sequence encoding a ZmRAD51 fusion protein, as well as the use of the plasmids to introduce the ZmRAD51 coding sequence into a host cell, such as maize cell, are also disclosed.

RE.CNT 3

RE.CNT

- (1) Smith, K; EMBL Sequence Data Library 1996
- (2) William, G; WO 9741228 A 1997 CAPLUS
- (3) Yeager Stassen, N; Current Genetics 1996
- => s l1 and (rice or barley or wheat)
- L6 344 L1 AND (RICE OR BARLEY OR WHEAT)
- => duplicate remove 16
- L7 184 DUPLICATE REMOVE L6 (160 DUPLICATES REMOVED)
- => s 17 and (gene or DNA)
- L8 184 L7 AND (GENE OR DNA)
- => d ti 1-50
- L8 ANSWER 1 OF 184 AGRICOLA
- TI Heterologous expression of genes mediating enhanced fungal resistance in transgenic wheat.
- L8 ANSWER 2 OF 184 AGRICOLA
- TI Regeneration of a lytic central vacuole and of neutral peripheral vacuoles can be visualized by green fluorescent proteins targeted to either type of vacuoles.
- L8 ANSWER 3 OF 184 AGRICOLA
- TI Enhanced resistance to sheath blight by constitutive expression of infection-related rice chitinase in transgenic elite indica rice cultivars.
- L8 ANSWER 4 OF 184 AGRICOLA
- TI **Chitinase** genes responsive to cold encode antifreeze proteins in winter cereals.
- L8 ANSWER 5 OF 184 AGRICOLA
- TI Expression of plant defence-related (PR-protein) transcripts during hardening and dehardening of winter wheat.
- L8 ANSWER 6 OF 184 AGRICOLA
- TI Flower-predominant expression of a  ${\tt gene}$  encoding a novel class I  ${\tt chitinase}$  in  ${\tt rice}$  (Oryza sativa L.).
- L8 ANSWER 7 OF 184 AGRICOLA
- TI Transgenic grapevine plants expressing a rice chitinase with enhanced resistance to fungal pathogens.
- L8 ANSWER 8 OF 184 AGRICOLA
- TI Fungal development and induction of defense response genes during early infection of wheat spikes by Fusarium graminearum.
- L8 ANSWER 9 OF 184 AGRICOLA
- TI Agrobacterium-mediated engineering for sheath blight resistance of indica rice cultivars from different ecosystems.
- L8 ANSWER 10 OF 184 AGRICOLA
- TI A transient assay system for the functional assessment of defense-related genes in wheat.
- L8 ANSWER 11 OF 184 AGRICOLA
- TI Transgenic chrysanthemum (Dendranthema grandiflorum (Ramat.) Kitamura) expressing a **rice chitinase gene** shows enhanced resistance to gray mold (Botrytis cinerea).
- L8 ANSWER 12 OF 184 AGRICOLA
- TI Enhanced resistance to blast (Magnaporthe grisea) in transgenic Japonica rice by constitutive expression of rice chitinase.
- L8 ANSWER 13 OF 184 AGRICOLA

- TI Elicitor action via cell membrane of a cultured **rice** cell demonstrated by the single-cell transient assay.
- L8 ANSWER 14 OF 184 AGRICOLA
- TI Genetic engineering of **wheat** for increased resistance to powdery mildew disease.
- L8 ANSWER 15 OF 184 AGRICOLA
- TI Development of **wheat** scab symptoms is delayed in transgenic **wheat** plants that constitutively express a **rice** thaumatin-like protein **gene**.
- L8 ANSWER 16 OF 184 AGRICOLA
- TI Regulation of the **chitinase gene** expression in suspension-cultured **rice** cells by N-acetylchitooligosaccharides: differences in the signal transduction pathways leading to the activation of elicitor-responsive genes.
- L8 ANSWER 17 OF 184 AGRICOLA
- TI Inheritance, expression, and silencing of a **chitinase** transgene in **rice**.
- L8 ANSWER 18 OF 184 AGRICOLA
- ${\tt TI}$  Candidate **gene** analysis of quantitative disease resistance in {\tt wheat.}
- L8 ANSWER 19 OF 184 AGRICOLA
- TI **Gene** activation of cytoplasmic acidification in suspension-cultured **rice** cells in response to the potent elicitor, N-acetylchitoheptaose.
- L8 ANSWER 20 OF 184 AGRICOLA
- TI Interactions between cold hardening and Microdochium nivale infection on expression of pathogenesis-related genes in winter wheat.
- L8 ANSWER 21 OF 184 AGRICOLA
- TI Introduction and constitutive expression of a rice chitinase gene in bread wheat using biolistic bombardment and the bar gene as a selectable marker.
- L8 ANSWER 22 OF 184 AGRICOLA
- TI Differential induction of apoplastic peroxidase and **chitinase** activities in susceptible and resistant **wheat** cultivars by Russian **wheat** aphid infestation.
- L8 ANSWER 23 OF 184 AGRICOLA
- TI Genetic engineering of disease resistance in cereals.
- L8 ANSWER 24 OF 184 AGRICOLA
- TI A new class II **rice chitinase**, Rcht2, whose induction by fungal elicitor is abolished by protein phosphatase 1 and 2A inhibitor.
- L8 ANSWER 25 OF 184 AGRICOLA
- TI Expression of a **chitinase** transgene in rose (Rosa hybrida L.) reduces development of blackspot disease (Diplocarpon rosae Wolf).
- L8 ANSWER 26 OF 184 AGRICOLA
- TI Transgenic cucumber plants harboring a **rice chitinase gene** exhibit enhanced resistance to gray mold (Botrytis cinerea).
- L8 ANSWER 27 OF 184 AGRICOLA
- TI Salicylic acid accumulation in **barley** is pathogen specific but not required for defense-**gene** activation.
- L8 ANSWER 28 OF 184 AGRICOLA
- TI Expression of beta-1,3-glucanase and **chitinase** in healthy, stem-rust-affected and elicitor-treated near-isogenic **wheat** lines showing Sr5- or Sr24-specified race-specific rust resistance.
- L8 ANSWER 29 OF 184 AGRICOLA
- TI Enhanced quantitative resistance against fungal disease by combinatorial expression of different **barley** antifungal proteins in transgenic tobacco.
- L8 ANSWER 30 OF 184 AGRICOLA
- TI Transgenic potato plants with enhanced resistance to the tomato moth, Lacanobia oleracea: growth room trials.
- L8 ANSWER 31 OF 184 AGRICOLA
- TI Regulation, expression and function of a new basic chitinase gene in rice (Oryza sativa L.).

L8 ANSWER 32 OF 184 AGRICOLA

- II Genetic engineering of rice for resistance to sheath blight.
- L8 ANSWER 33 OF 184 AGRICOLA
- TI Acquired resistance in **barley**. The resistance mechanisms induced by 2,6-dichloroisonicotinic acid is a phenocopy of a genetically based mechanism governing race-specific powdery mildew resistance.
- L8 ANSWER 34 OF 184 AGRICOLA
- TI Identification of an enhancer/silencer sequence directing the aleurone-specific expression of a barley chitinase gene.
- L8 ANSWER 35 OF 184 AGRICOLA
- TI Molecular and cellular expression of quantitative resistance in **barley** to powdery mildew.
- L8 ANSWER 36 OF 184 AGRICOLA
- TI Nar-1 and Nar-2, two loci required for Mlal2-specified race-specific resistance to powdery mildew in **barley**.
- L8 ANSWER 37 OF 184 AGRICOLA
- TI The relationship between the expression of defense-related genes and mildew development in **barley**.
- L8 ANSWER 38 OF 184 AGRICOLA
- TI Sequence variation, differential expression and chromosomal location of rice chitinase genes.
- L8 ANSWER 39 OF 184 AGRICOLA
- TI Stress induction and developmental regulation of a rice chitinase promoter in transgenic tobacco.
- L8 ANSWER 40 OF 184 AGRICOLA
- TI Molecular basis of post-translational modifications and targeting of **barley** lectin to the vacuoles in **barley** and in transgenic tobacco plants.
- L8 ANSWER 41 OF 184 AGRICOLA
- TI The **barley** lectin carboxyl-terminal propeptide is a vacuolar protein sorting determinant in plants.
- L8 ANSWER 42 OF 184 AGRICOLA
- TI Identification of an endochitinase cDNA clone from **barley** aleurone cells.
- L8 ANSWER 43 OF 184 AGRICOLA
- TI Isolation and characterization of a **rice gene** encoding a basic **chitinase**.
- L8 ANSWER 44 OF 184 AGRICOLA
- TI cDNA cloning and characterization of a putative 1,3-beta-glucanase transcript induced by fungal elicitor in bean cell suspension cultures.
- L8 ANSWER 45 OF 184 AGRICOLA
- TI Biochemical and molecular characterization of three **barley** seed proteins with antifungal properties.
- L8 ANSWER 46 OF 184 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Stylosanthes guianensis plants transformed with a **rice chitinase gene** confer resistance to Rhizoctonia foliar blight disease.
- L8 ANSWER 47 OF 184 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Use of cyanamide hydratase **gene** as a selectable marker for the transformation of sorghum.
- L8 ANSWER 48 OF 184 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Transgenic wheat plants: A powerful breeding source.
- L8 ANSWER 49 OF 184 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Complementary DNA for rice chitinase having lytic activity against moulds and bacteria, and vector containing said complementary DNA and transformant.
- L8 ANSWER 50 OF 184 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Isolation and characterization of novel cDNA clones of acidic chitinases and beta-1,3-glucanases from **wheat** spikes infected by Fusarium graminearum.

- L7 ANSWER 1 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Transgenic tobacco expressing NtmybPS1 and PA1 genes for pathogen resistance
- L7 ANSWER 2 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Cloning of a vascular-specific promoter from rice Oshox1
  gene and its use in making transgenic plants
- L7 ANSWER 3 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Nucleic acid sequences encoding cell wall-degrading enzymes and their use to engineer plant resistance to Fusarium and other pathogens
- L7 ANSWER 4 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI DNA constructs encoding chimeric plant RRK receptors (Bril::Xa21 and Hevein::Xa21), and their use in production of transgenic plants
- L7 ANSWER 5 OF 184 AGRICOLA DUPLICATE 1
- TI Heterologous expression of genes mediating enhanced fungal resistance in transgenic wheat.
- L7 ANSWER 6 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Hydrophobic interactions of the structural protein GRP1.8 in the cell wall of protoxylem elements
- L7 ANSWER 7 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Construction of plant expression vector containing multiple gene
- L7 ANSWER 8 OF 184 AGRICOLA DUPLICA
- TI Enhanced resistance to sheath blight by constitutive expression of infection-related **rice chitinase** in transgenic elite indica **rice** cultivars.
- L7 ANSWER 9 OF 184 CABA COPYRIGHT 2001 CABI
- TI Enhanced resistance to sheath blight by constitutive expression of infection-related **rice chitinase** in transgenic elite indica **rice** cultivars.
- L7 ANSWER 10 OF 184 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 3
- TI Isolation and characterization of novel cDNA clones of acidic chitinases and beta-1,3-glucanases from **wheat** spikes infected by Fusarium graminearum.
- L7 ANSWER 11 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Introduction of rice chitinase gene into wheat via low energy Ar+ beam implantation
- L7 ANSWER 12 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI A Survey of the Molecular Evolutionary Dynamics of Twenty-Five Multigene Families from Four Grass Taxa
- L7 ANSWER 13 OF 184 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 4
- TI Transgenic wheat plants: A powerful breeding source.
- L7 ANSWER 14 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Novel transgenic **rice** strains resistant to blast and sheathblight
- L7 ANSWER 15 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Rapid development of homozygous transgenic **rice** using anther culture harboring **rice chitinase gene** for enhanced sheath blight resistance
- L7 ANSWER 16 OF 184 AGRICOLA
- TI Regeneration of a lytic central vacuole and of neutral peripheral vacuoles can be visualized by green fluorescent proteins targeted to either type of vacuoles.
- L7 ANSWER 17 OF 184 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Stylosanthes guianensis plants transformed with a **rice chitinase gene** confer resistance to Rhizoctonia foliar blight disease.
- L7 ANSWER 18 OF 184 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Use of cyanamide hydratase **gene** as a selectable marker for the transformation of sorghum.
- L7 ANSWER 19 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Transformation of roses with genes for antifungal proteins
- L7 ANSWER 20 OF 184 CABA COPYRIGHT 2001 CABI

- TI Transformation of roses with genes for antifungal proteins.
- L7 ANSWER 21 OF 184 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Complementary **DNA** for **rice chitinase** having lytic activity against moulds and bacteria, and vector containing said complementary **DNA** and transformant.
- L7 ANSWER 22 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Method for producing transgenic plants resistant to glyphosate herbicides
- L7 ANSWER 23 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Method for producing transgenic plants resistant to glyphosate herbicides
- L7 ANSWER 24 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Method for producing transgenic plants resistant to glyphosate herbicides
- L7 ANSWER 25 OF 184 CAPLUS COPYRIGHT 2001 ACS
- $\mbox{TI}\mbox{ }\mbox{ Maize chitinases and cDNAs and method of modulating $\mbox{\it chitinase}$ activity in plants }$
- L7 ANSWER 26 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Inducible plant caffeic acid O-methyltransferase II **gene** promoter and chimeric genes for expression in plants
- L7 ANSWER 27 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Protein and cDNA sequences of Urtica dioica agglutinin and uses thereof in plant disease control
- L7 ANSWER 28 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Wheat chitinase genes expressed at low temperature and method for its isolation
- L7 ANSWER 29 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Phytopathogenic fungi control agents containing chitinase
- L7 ANSWER 30 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Preparation of transgenic grapes expressing **chitinase** for fungal disease resistance
- L7 ANSWER 31 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Pathogen inducible promoter from Arabidopsis thaliana and its use in preparation of pathogen-resistance transgenic plants
- L7 ANSWER 32 OF 184 AGRICOLA DUPLICATE 5
- TI **Chitinase** genes responsive to cold encode antifreeze proteins in winter cereals.
- L7 ANSWER 33 OF 184 AGRICOLA DUPLICATE
- TI Flower-predominant expression of a **gene** encoding a novel class I **chitinase** in **rice** (Oryza sativa L.).
- L7 ANSWER 34 OF 184 AGRICOLA DUPLICATE 7
- ${\tt TI}$  Agrobacterium-mediated engineering for sheath blight resistance of indica rice cultivars from different ecosystems.
- L7 ANSWER 35 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Cerebroside elicitors found in diverse phytopathogens activate defense responses in **rice** plants
- L7 ANSWER 36 OF 184 AGRICOLA DUPLICATE 8
- TI Transgenic grapevine plants expressing a rice chitinase with enhanced resistance to fungal pathogens.
- L7 ANSWER 37 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Transformation system and study on the introduction of  ${\bf chitinase}$   ${\bf gene}$  into Linum usitatisium
- L7 ANSWER 38 OF 184 AGRICOLA DUPLICATE 9
- TI Elicitor action via cell membrane of a cultured **rice** cell demonstrated by the single-cell transient assay.
- L7 ANSWER 39 OF 184 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 10
- TI Isolation and expression of a pistil-specific chitinase gene in rice (Oryza sativa L.
- L7 ANSWER 40 OF 184 AGRICOLA DUPLICATE 11
- TI Fungal development and induction of defense response genes during early infection of wheat spikes by Fusarium graminearum.
- L7 ANSWER 41 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Identification of **chitinase** and .beta.-1,3-glucanase cDNAs from citrus fruit

- L7 ANSWER 42 OF 184 CABA COPYRIGHT 2001 CABI
- TI Identification of **chitinase** and beta -1,3-glucanase cDNAs from citrus fruit.
- L7 ANSWER 43 OF 184 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 12
- TI Influence of fungal-bacterial interactions on bacterial conjugation in the residuesphere.
- L7 ANSWER 44 OF 184 CABA COPYRIGHT 2001 CABI
- TI Regeneration of transgenic tomato plants expressing **rice chitinase gene** via Agrobacterium tumefaciens-mediated transformation.
- L7 ANSWER 45 OF 184 AGRICOLA

- TI Expression of plant defence-related (PR-protein) transcripts during hardening and dehardening of winter **wheat**.
- L7 ANSWER 46 OF 184 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 14
- TI Transgenic expression of cecropin B, an antibacterial peptide from Bombyx mori, confers enhanced resistance to bacterial leaf blight in **rice**
- L7 ANSWER 47 OF 184 CAPLUS COPYRIGHT 2001 ACS DUPLICATE 15
- TI Transgenic tobacco plant expressing a **rice chitinase** and their resistance to Alternaria alternata
- L7 ANSWER 48 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Transgene identification in transgenic seeds using screenable markers linked to aleurone-specific promoters
- L7 ANSWER 49 OF 184 CAPLUS COPYRIGHT 2001 ACS
- TI Genes for enzymes of salicylate biosynthesis of for the induction of disease resistance in plants
- L7 ANSWER 50 OF 184 CAPLUS COPYRIGHT 2001 ACS
- ${\tt TI}$  Cloning expression systems and cDNA sequences encoding antifreeze proteins from winter rye
- => s 14 and (rice or barley or wheat)
- L9 4 L4 AND (RICE OR BARLEY OR WHEAT)
- => d ti 1-4
- L9 ANSWER 1 OF 4 CAPLUS COPYRIGHT 2001 ACS
- TI Homologous recombination and molecular evolution of recombination protein homologs in plants
- L9 ANSWER 2 OF 4 CAPLUS COPYRIGHT 2001 ACS
- TI Gene RAD51 recombinase and cDNA of corn and wheat and use of RAD51 in positive selection of recombinant plant cells
- L9 ANSWER 3 OF 4 CAPLUS COPYRIGHT 2001 ACS
- TI Protein and cDNA sequences of maize RAD51-like gene and uses thereof in transgenic plants
- L9 ANSWER 4 OF 4 CAPLUS COPYRIGHT 2001 ACS
- TI Identification of a novel human RAD51 homolog, RAD51B. [Erratum to document cited in CA128:201583]
- => d bib abs 2 3 1
- L9 ANSWER 2 OF 4 CAPLUS COPYRIGHT 2001 ACS
- AN 2000:814630 CAPLUS
- DN 133:360459
- TI Gene RAD51 recombinase and cDNA of corn and wheat and use of RAD51 in positive selection of recombinant plant cells

A2 20001116

- IN Famodu, Omolayo O.; Morgante, Michele
- PA E.I. du Pont de Nemours and Company, USA
- SO PCT Int. Appl., 46 pp. CODEN: PIXXD2

WO 2000068390

- DT Patent
- LA English
- FAN. CNT 1

PT

PATENT NO. KIND DATE

APPLICATION NO. DATE

WO 2000-US12587 20000509

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20010503
     WO 2000068390
                         АЗ
          W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU,
              CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL,
              IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD,
              MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK,
              SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZW, AM, AZ, BY, KG,
              KZ, MD, RU, TJ, TM
          RW: GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE,
              DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
PRAI US 1999-133438
                       P
                             19990511
     This invention relates to cDNAs encoding corn and wheat gene
     RAD51 recombinases. The invention also relates to the
     construction of a chimeric gene encoding all or a substantial portion of
     the recombination protein, in sense or antisense orientation, wherein
     expression of the chimeric gene results in prodn. of altered levels of the
     recombination protein in a transformed host cell. Also disclosed is use
     of the chimeric gene for pos. selection of transformed plant cells.
1.9
     ANSWER 3 OF 4 CAPLUS COPYRIGHT 2001 ACS
     2000:814610 CAPLUS
AN
     134:1353
DN
ΤI
     Protein and cDNA sequences of maize RAD51-like gene and uses
     thereof in transgenic plants
IN
     Mahajan, Pramod B.; Shi, Jinrui
     Pioneer Hi-Bred International, Inc., USA
PA
SO
     PCT Int. Appl., 76 pp.
     CODEN: PIXXD2
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     PATENT NO.
                       KIND DATE
                                               APPLICATION NO. DATE
                        ____
     WO 2000068370 A2
WO 2000068370 A3
                               20001116
                                               WO 2000-US9010 20000405
РΤ
                               20010215
          W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU,
              CZ, DE, DK, DM, EE, ES, FI, GB, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD,
              MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK,
              SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, AM, AZ,
              BY, KG, KZ, MD, RU, TJ, TM
          RW: GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF,
              CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
PRAI US 1999-132582
                       P
                             19990505
     The invention provides protein and cDNA sequences of a novel maize
     ortholog of RAD51, which shows high homol. to the human RAD51C
     gene. The present invention provides methods and compns. relating to
     altering maize RAD51 levels in plants. The invention further
     provides recombinant expression cassettes, host cells, transgenic plants,
     and antibody compns.
L9
     ANSWER 1 OF 4 CAPLUS COPYRIGHT 2001 ACS
     2001:397024 CAPLUS
ΑN
DN
     135:1212
TI
     Homologous recombination and molecular evolution of recombination protein
     homologs in plants
TN
     Lassner, Michael; Delcardayre, Steven
PA
     Maxygen, Inc., USA
     PCT Int. Appl., 57 pp.
     CODEN: PIXXD2
DT
     Patent
LA
     English
FAN.CNT 1
     PATENT NO.
                        KIND DATE
                                                APPLICATION NO.
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     WO 2001038504
PI
                         A2
                               20010531
                                                WO 2000-US32289
                                                                   20001122
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              LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU,
              SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
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              BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
-167450 P 19991123
PRAI US 1999-167450
   Methods for evolving recombinase protein homologs and RecA/VirE2 fusion
     proteins which complement VirE2 deficient Agrobacterium are provided. The
     use of recombinase protein homologs and RecA/VirE2 fusion proteins in the
     context of Agrobacterium-mediated transformation are provided. Methods
     for producing transgenic organisms by homologous recombination using
```

evolved recombinase proteins and Agrobacterium strains which express recombinase protein homologs or RecA/VirE2 fusion proteins are provided. Transgenic cells and organisms which have integrated an exogenous DNA sequence into a predetd. site in their genome are provided.

=> logoff hold

FILE 'HOME' ENTERED AT 08:24:16 ON 22 OCT 2001

- => file agricola biosis caplus caba
- => s rad5? and (plant or maize or wheat or tobacco or arabidopsis or rice or sunflower)
- L1 83 RAD5? AND (PLANT OR MAIZE OR WHEAT OR TOBACCO OR ARABIDOPSIS OR RICE OR SUNFLOWER)
- => duplicate remove 11
- L2 49 DUPLICATE REMOVE L1 (34 DUPLICATES REMOVED)
- => d ti 1-49
- L2 ANSWER 1 OF 49 CAPLUS COPYRIGHT 2001 ACS
- ${\tt TI}$  Methods of improving homologous recombination for gene targeting and therapy
- L2 ANSWER 2 OF 49 CAPLUS COPYRIGHT 2001 ACS
- ${
  m TI}$  Homologous recombination and molecular evolution of recombination protein homologs in plants
- L2 ANSWER 3 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI Methods of determining individual hypersensitivity to a pharmaceutical agent from gene expression profile
- L2 ANSWER 4 OF 49 CABA COPYRIGHT 2001 CABI
- TI A RecA homologue in Ustilago maydis that is distinct and evolutionarily distant from **Rad51** actively promotes DNA pairing reactions in the absence of auxiliary factors.
- L2 ANSWER 5 OF 49 AGRICOLA

DUPLICATE 1

- TI RAD50 function is essential for telomere maintenance in Arabidopsis.
- L2 ANSWER 6 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 2
- TI In vitro and in vivo nucleotide exchange directed by chimeric RNA/DNA oligonucleotides in Saccharomyces cerevisae.
- L2 ANSWER 7 OF 49 CAPLUS COPYRIGHT 2001 ACS DUPLICATE 3
- ${\tt TI}$  Homologous recombination in planta is stimulated in the absence of  ${\tt Rad50}$
- L2 ANSWER 8 OF 49 AGRICOLA

- TI Disruption of the **Arabidopsis RAD50** gene leads to **plant** sterility and MMS sensitivity.
- L2 ANSWER 9 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI Protein and cDNA sequences of maize RAD50-like gene and uses thereof in transgenic plants
- L2 ANSWER 10 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI Gene RAD51 recombinase and cDNA of corn and wheat and use of RAD51 in positive selection of recombinant plant cells
- L2 ANSWER 11 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI Protein and cDNA sequences of maize RAD51-like gene and uses thereof in transgenic plants
- L2 ANSWER 12 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI Locked nucleic acid hybrids and their uses in modulating genetic processes
- L2 ANSWER 13 OF 49 CAPLUS COPYRIGHT 2001 ACS
- ${
  m TI}$  Small protein SML1 that interacts with yeast ribonucleotide reductase subunit
- L2 ANSWER 14 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS
- TI The FHA domain mediates phosphoprotein interactions.
- L2 ANSWER 15 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 5
- TI MUS81 encodes a novel helix-hairpin-helix protein involved in the response

to UV- and methylation-induced DNA damage in Saccharomyces cerevisiae.

- L2 ANSWER 16 OF 49 CABA COPYRIGHT 2001 CABI
- TI Isolation and biochemical characterization of a new topoisomerase I inhibitor from Ocotea leucoxylon.
- L2 ANSWER 17 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 6
- TI Elevated UV-B radiation reduces genome stability in plants.
- L2 ANSWER 18 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 7
- TI The bacterial replicative helicase DnaB evolved from a RecA duplication.
- L2 ANSWER 19 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI Nucleotide sequences encoding maize RAD51 recombinases
- L2 ANSWER 20 OF 49 AGRICOLA

DUPLICATE

- TI Three-dimensional microscopy of the Rad51 recombination protein during meiotic prophase.
- L2 ANSWER 21 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS
- TI DNA repair in lymphoblastoid cell lines from patients with head and neck cancer.
- L2 ANSWER 22 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI **Plant** responses to genotoxic stress are linked to an ABA/salinity signaling pathway
- L2 ANSWER 23 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI Green fluorescent protein and yeast DNA repair genes in construction of recombinant DNA mol. for detection of DNA-damaging agents
- L2 ANSWER 24 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 9
- TI The human REC2/RAD51B gene acts as a DNA damage sensor by inducing G1 delay and hypersensitivity to ultraviolet irradiation.
- L2 ANSWER 25 OF 49 AGRICOLA

DUPLICATE 1

DUPLICATE 12

- TI Agrobacterium tumefaciens transformation of the radiation hypersensitive **Arabidopsis** thaliana mutants uvhl and **rad5**.
- L2 ANSWER 26 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI Identification of a novel human RAD51 homolog, RAD51B.
  [Erratum to document cited in CA128:201583]
- L2 ANSWER 27 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Rad51 distribution is altered in desynaptic2, a maize meiotic mutant that has abnormal chromosome pairing.
- L2 ANSWER 28 OF 49 AGRICOLA DUPLICATE 11
- TI Isolation and characterisation of the  ${\bf RAD51}$  and DMC1 homologs from  ${\bf Arabidopsis}$  thaliana.
- L2 ANSWER 29 OF 49 AGRICOLA
- TI **Arabidopsis** thaliana mutants altered in homologous recombination.
- L2 ANSWER 30 OF 49 CABA COPYRIGHT 2001 CABI
- TI Interaction between Ustilago maydis REC2 and RAD51 genes in DNA repair and mitotic recombination.
- L2 ANSWER 31 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 13
- TI Isolation and characterization of **rad51** orthologs from Coprinus cinereus and Lycopersicon esculentum, and phylogenetic analysis of eukaryotic recA homologs.
- L2 ANSWER 32 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Cap-prevented recombination between terminal telomeric repeat arrays (telomere CPR) maintains telomeres in Kluyveromyces lactis lacking telomerase.
- L2 ANSWER 33 OF 49 AGRICOLA

- TI Nucleotide sequences of nuclear tRNA(Cys) genes from Nicotiana and **Arabidopsis** and expression in HeLa cell extract.
- L2 ANSWER 34 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS
- ${\tt TI}$  Using 3-D immunocytology to explore the roles of mismatch binding proteins in meiosis.
- L2 ANSWER 35 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI Identification and expression of the Neurospora crassa mei-3 gene which encodes a protein homologous to **Rad51** of Saccharomyces cerevisiae

- L2 ANSWER 36 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI Characterization of a mouse recA-like gene specifically expressed in testis
- L2 ANSWER 37 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI Structural analysis of a recA-like gene in the genome of **Arabidopsis** thaliana
- L2 ANSWER 38 OF 49 CABA COPYRIGHT 2001 CABI
- TI Structure of REC2, a recombinational repair gene of Ustilago maydis, and its function in homologous recombination between plasmid and chromosomal sequences.
- L2 ANSWER 39 OF 49 AGRICOLA

- TI Isolation of **Arabidopsis** thaliana mutants hypersensitive to gamma radiation.
- L2 ANSWER 40 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI Characterization of cDNAs induced in meiotic prophase in lily microsporocytes
- L2 ANSWER 41 OF 49 CAPLUS COPYRIGHT 2001 ACS
- TI Stabilization of a yeast artificial chromosome containing **plant**DNA using a recombination-deficient host
- L2 ANSWER 42 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Expression of the Saccharomyces cerevisiae PAD50 gene during meiosis: Steady-state transcript levels rise and fall while steady-state protein levels remain constant.
- L2 ANSWER 43 OF 49 AGRICOLA

DUPLICATE 16

- TI Isolation and characterization of a yeast gene that is homologous with a meiosis-specific cDNA from a **plant**.
- L2 ANSWER 44 OF 49 CABA COPYRIGHT 2001 CABI
- TI The use of saccharomycete yeasts to evaluate genetic effects of systemic fungicides.
- L2 ANSWER 45 OF 49 AGRICOLA
- TI A Saccharomyces cerevisiae RAD52 allele expressing a C-terminal truncation protein: activities and intragenic complementation of missense mutations.
- L2 ANSWER 46 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS
- TI Characterization of REC104, a gene required for early meiotic recombination in the yeast Saccharomyces cerevisiae.
- L2 ANSWER 47 OF 49 BIOSIS COPYRIGHT 2001 BIOSIS
- TI THE DNA REPAIR GENE PSO3 OF SACCHAROMYCES-CEREVISIAE BELONGS TO THE RAD3 EPISTASIS GROUP.
- L2 ANSWER 48 OF 49 AGRICOLA
- TI The rec102 mutant of yeast is defective in meiotic recombination and chromosome synapsis.
- L2 ANSWER 49 OF 49 AGRICOLA
- TI Nucleotide sequence of the RAD57 gene of Saccharomyces cerevisiae.
- => d bib abs 43 37 28 19 22 7-10 2 5
- L2 ANSWER 43 OF 49 AGRICOLA

DUPLICATE 16

AN 93:47656 AGRICOLA

DN IND93029549

- TI Isolation and characterization of a yeast gene that is homologous with a meiosis-specific cDNA from a **plant**.
- AU Kobayashi, T.; Hotta, Y.; Tabata, S.
- CS Nagoya University, Nagoya, Japan
- AV DNAL (442.8 Z34)
- SO M G G: Molecular and general genetics, Feb 1993. Vol. 237, No. 1/2. p. 225-232

Publisher: Berlin, W. Ger. : Springer International.

CODEN: MGGEAE; ISSN: 0026-8925 NTE Includes references.

- DT Article
- FS Non-U.S. Imprint other than FAO
- LA English
- AB By using as probe a meiosis-specific cDNA clone LIM15 from the monocotyledonous **plant**, Lilium longiflorum, a clone containing a 2.8 kb DNA fragment was isolated from a genomic library of Saccharomyces

cerevisiae. Primary structure analysis revealed that the clone includes two complete open reading frames, designated ISC2 and ISC10, capable of coding for a 36.6 kDa and a 31.6 kDa polypeptide, respectively, with the former frame being interrupted by a 92 bp intron. The predicted amino acid sequence of Isc2 was 56% identical with the putative gene product of lily cDNA clone LIM15, and showed limited sequence similarity with the yeast RAD57 gene product. Transcripts of the two genes begin accumulating 2.5 h and 7.5 h after induction of meiosis, respectively, according to a Northern hybridization analysis. Since disruption of either one of these genes had a drastic effect on the ability to form spores, ISC2 and ISC10 are expected to play significant roles in the formation of

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reproductive cells.
     ANSWER 37 OF 49 CAPLUS COPYRIGHT 2001 ACS
L_2
     1995:631987 CAPLUS
AN
DN
     123:221650
     Structural analysis of a recA-like gene in the genome of
     Arabidopsis thaliana
     Sato, Shusei; Hotta, Yasuo; Tabata, Satoshi
ΑH
CS
     Kazusa DNA Res. Inst., Chiba, 292, Japan
SO
     DNA Res. (1995), 2(2), 89-93
     CODEN: DARSE8; ISSN: 1340-2838
DТ
     Journal
LA
     English
     A recA-like gene was identified in the genome of Arabidopsis
     thaliana by means of PCR using primers designed on the basis of previously
     reported amino acid sequences of eukaryotic RecA-like proteins. The
     structure of the gene, termed ArLIM15, was investigated by comparing the
     primary structure of the genomic DNA with that of the corresponding cDNA.
     The open reading frame, which was split into 15 exons, was established to
     have the capacity for encoding a 37.3-kDa polypeptide. The amino acid
     sequence of the putative product of ArLIM15 showed a high degree of
     similarity to that of LIM15 in the monocotyledonous plant
     Lilium, including a 93% identity, and to those of other recA-like genes in
     yeasts and vertebrates with identities of 69-71%. Phylogenetic anal.
     indicated ArLIM15 to be much closer to meiosis-specific LIM15 and DMC1 in
     Saccharomyces cerevisiae than to RAD51 in S. cerevisiae and its
     homologs on an evolutionary scale.
L2
     ANSWER 28 OF 49 AGRICOLA
                                                            DUPLICATE 11
     1998:62619 AGRICOLA
AN
DN
     IND21241600
     Isolation and characterisation of the {\it RAD51} and DMC1 homologs
TΙ
     from Arabidopsis thaliana.
ΑU
     Doutriaux, M.P.; Couteau, F.; Bergounioux, C.; White, C.
ΑV
     DNAL (442.8 Z34)
     Molecular & general genetics : MGG, Feb 1998. Vol. 257, No. 3. p. 283-291
SO
     Publisher: Berlin, Germany: Springer-Verlag Berlin.
     CODEN: MGGEAE; ISSN: 0026-8925
NTE
     Includes references
CY
     Germany
DT
     Article
FS
     Non-U.S. Imprint other than FAO
LA
     English
     ANSWER 19 OF 49 CAPLUS COPYRIGHT 2001 ACS
     1999:529275 CAPLUS
AN
DN
     131:154494
     Nucleotide sequences encoding {\tt maize} RAD51 recombinases
TΙ
ΙN
     Bowen, Benjamin A.; Chamberlin, Mark A.; Drummond, Bruce J.; Mcelver, John
     A.; Rothstein, Rodney J.
PΑ
     Pioneer Hi-Bred International, Inc., USA
SO
     PCT Int. Appl., 66 pp.
     CODEN: PIXXD2
DT
      Patent
     English
LA
FAN.CNT 1
      PATENT NO.
                        KIND DATE
                                              APPLICATION NO. DATE
PΙ
     WO 9941394
                        A1 19990819
                                              WO 1999-US2900
                                                                19990211
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              KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW,
              MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, UZ, VN, YU, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
          RW: GH, GM, KE, LS, MW, SD, SZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES,
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A1 19990830 A1 20001122

20001122

A1

AU 9926699 EP 1053339

IE. FI

AU 1999-26699

R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,

EP 1999-906894

19990211

19990211

Р 19980213 PRAI US 1998-74745 WO 1999-US2900 W 19990211 Nucleic acid sequences encoding two RAD51 recombinases active in maize plants are provided. Full-length cDNA sequences from a maize tassel cDNA library, including the ZmRAD51 coding sequences and unique 3-untranslated regions which are useful as RFLP probes, are also provided. RFLP mapping indicates that the Zea mays genome contains 2 genes encoding different variants of the ZmRAD51 recombinase enzyme, one mapped to chromosome 7 and the other on chromosome 3. The prodn. of plasmids contg. a nucleic acid sequence encoding a ZmRAD51 fusion protein, as well as the use of the plasmids to introduce the ZmRAD51 coding sequence into a host cell, such as maize cell, are also disclosed. RE.CNT 3 RE (1) Smith, K; EMBL Sequence Data Library 1996 (2) William, G; WO 9741228 A 1997 CAPLUS (3) Yeager Stassen, N; Current Genetics 1996 L2 ANSWER 22 OF 49 CAPLUS COPYRIGHT 2001 ACS 1999:141888 CAPLUS ΑN DN 130:309109 ΤI Plant responses to genotoxic stress are linked to an ABA/salinity signaling pathway Albinsky, Doris; Masson, Jean E.; Bogucki, Augustyn; Afsar, Karin; Vass, ΑIJ Imre; Nagy, Ferenc; Paszkowski, Jerzy Friedrich Miescher-Institute, Basel, CH-4002, Switz. CS SO Plant J. (1999), 17(1), 73-82 CODEN: PLJUED: ISSN: 0960-7412 PR Blackwell Science Ltd. DT Journal LA Enalish AR An Arabidopsis mutant (uvs66) is described that appears to be affected in the perception of signals triggered by genotoxic treatments. The mutant uvs66 was identified as hypersensitive to UV-C and to the  ${\sf DNA-damaging}$  chems. Me methane sulfonate (MMS) and mitomycin C (MMC), but seems to perform light dependent repair, nucleotide excision repair and homologous recombinational repair as efficiently as the wild type. Exposure of uvs66 plants to various environmental stresses revealed a normal response, with the exception of elevated salinity and abscisic acid (ABA). The hypersensitivity to NaCl and ABA is correlated with aberrant regulation of transcripts that are regulated by ABA (RAB18), or are induced by DNA damaging treatments (AtRAD51). The properties of the mutant uvs66 suggest an unexpected link between ABA and/or salt stress mediated signals and genotoxic stress responses, and provide an important connection between the physiol. and genetic responses of plants to abiotic stress factors. RE.CNT 53 RE (1) Aro, E; Biochem Biophys Acta 1993, V1143, P113 CAPLUS (2) Basile, G; Mol Cell Biol 1992, V12, P3235 CAPLUS (3) Baur, M; Mol Cell Biol 1990, V10, P492 CAPLUS (5) Bohnert, H; Plant Cell 1995, V7, P1099 CAPLUS (6) Bostock, R; Plant Physiol 1992, V98, P1356 CAPLUS ALL CITATIONS AVAILABLE IN THE RE FORMAT ANSWER 7 OF 49 CAPLUS COPYRIGHT 2001 ACS 2001:464972 CAPLUS L2 DUPLICATE 3 AN ΤI Homologous recombination in planta is stimulated in the absence of Rad50 ΑU Gherbi, Hassen; Gallego, Maria Eugenia; Jalut, Nicole; Lucht, Jan M.; Hohn, Barbara; White, Charles I. CS UMR 6547 BIOMOVE, Universite Blaise Pascal, Aubiere, 63177, Fr. SO EMBO Rep. (2001), 2(4), 287-291 CODEN: ERMEAX; ISSN: 1469-221X PR Oxford University Press DΤ Journal LA English AB Chromosomal double-strand DNA breaks must be repaired; in the absence of repair the resulting acentrometric (and telomereless) fragments may be lost and/or the broken DNA ends may recombine causing general chromosomal instability. The Rad50/Mrell/Xrs2 protein complex acts at DNA ends and is implicated in both homologous and non-homologous recombination. We have isolated a rad50 mutant of the plant Arabidopsis thaliana and show here that it has a somatic hyper-recombination phenotype in planta. This finding supports the hypothesis of a competition between homologous and illegitimate

recombination in higher eukaryotes. To our knowledge, this is the first direct in vivo support for the role of this complex in chromosomal recombination in a multicellular organism and the first description of a

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(1) Berthold, D; Plant Mol Biol Rep 1993, V11, P338 CAPLUS
(4) Critchlow, S; Trends Biochem Sci 1998, V23, P394 CAPLUS
(5) Gallego, M; Plant J 2001, V25, P31 CAPLUS
(7) Gottlieb, S; Proc Natl Acad Sci USA 1989, V86, P7072 CAPLUS (8) Haber, J; Nature 1999, V398, P665 CAPLUS
ALL CITATIONS AVAILABLE IN THE RE FORMAT
                                                          DUPLICATE 4
     ANSWER 8 OF 49 AGRICOLA
     2001:37560 AGRICOLA
ΑN
DN
     TND22438193
     Disruption of the Arabidopsis RAD50 gene leads to
     plant sterility and MMS sensitivity.
     Gallego, M.E.; Jeanneau, M.; Granier, F.; Bouchez, D.; Bechtold, N.;
AU
     White, C.I.
     DNAL (QK710.P68)
ΑV
     The Plant journal: for cell and molecular biology, Jan 2001. Vol. 25, No.
SO
     1. p. 31-41
     Publisher: Oxford : Blackwell Sciences Ltd.
     ISSN: 0960-7412
NTE
     Includes references
     England; United Kingdom
CY
DT
     Article
     Non-U.S. Imprint other than FAO
FS
LA
     English
     The Rad50 protein is involved in the cellular response to
AB
     DNA-double strand breaks (DSBs), including the detection of damage,
     activation of cell-cycle checkpoints, and DSB repair via recombination. It
     is essential for meiosis in yeast, is involved in telomere maintenance,
     and is essential for cellular viability in mice. Here we present the
     isolation, sequence and characterization of the Arabidopsis
     thaliana RAD50 homologue (AtRAD50) and an Arabidopsis
     mutant of this gene. A single copy of this gene is present in the
     Arabidopsis genome, located on chromosome II. Northern analysis
     shows a single 4.3 Kb mRNA species in all plant tissues tested,
     which is strongly enriched in flowers and other tissues with many dividing
     cells. The predicted protein presents strong conservation with the other
     known {\bf Rad50} homologues of the amino- and carboxy-terminal
     regions. Mutant plants present a sterility phenotype which co-segregates
     with the T-DNA insertion. Molecular analysis of the mutant plants shows
     that the sterility phenotype is present only in the plants homozygous for
     the T-DNA insertion. An in vitro mutant cell line, derived from the mutant
     plant, shows a clear hypersensitivity to the DNA-damaging agent
     methylmethane sulphonate, suggesting a role of RAD50 in
     double-strand break repair in plant cells. This is the first
     report of a {\tt plant} mutated in a protein of the {\tt Rad50}
     -Mrell-Xrs2 complex, as well as the first data suggesting the involvement
     of the Rad50 homologue protein in meiosis and DNA repair in
     plants.
     ANSWER 9 OF 49 CAPLUS COPYRIGHT 2001 ACS
L2
     2000:814642 CAPLUS
ΑN
DN
     134:1355
     Protein and cDNA sequences of maize RAD50-like gene
TТ
     and uses thereof in transgenic plants
IN
     Mahajan, Pramod B.; Shi, Jinrui
     Pioneer Hi-Bred International, Inc., USA
PA
     PCT Int. Appl., 78 pp.
SO
     CODEN: PIXXD2
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LA
     English
FAN. CNT 1
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     WO 2000068404
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PΙ
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     EP 1093523
                       A1 20010425
                                             EP 2000-931950 20000425
         R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,
             IE, SI, LT, LV, FI, RO
PRAI US 1999-132575
                       Р
                             19990505
     WO 2000-US11086 W
                             20000425
     The invention provides protein and cDNA sequences of a novel maize
     ortholog of RAD50, which shows high homol. to the yeast
     RAD50 gene. The present invention provides methods and compns.
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relating to altering maize RAD50 levels in plants.
     The invention further provides recombinant expression cassettes, host
     cells, transgenic plants, and antibody compns.
RE.CNT 2
RE
   Genelabs Tech Inc; WO 9838306 A 1998 CAPLUS
(1)
(2) Petrini, J; American Journal of Human Genetics 1999, V64, P1264 CAPLUS
     ANSWER 10 OF 49 CAPLUS COPYRIGHT 2001 ACS
1.2
ΑN
     2000:814630 CAPLUS
DN
     133:360459
     Gene RAD51 recombinase and cDNA of corn and wheat and
TΙ
     use of RAD51 in positive selection of recombinant plant
TN
     Famodu, Omolayo O.; Morgante, Michele
     E.I. du Pont de Nemours and Company, USA
PA
SO
     PCT Int. Appl., 46 pp.
     CODEN: PIXXD2
     Patent
LA
     English
FAN.CNT 1
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                                           APPLICATION NO. DATE
                      A2
     WO 2000068390
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                                           WO 2000-US12587 20000509
PΙ
                            20010503
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             KZ, MD, RU, TJ, TM
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             DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF,
             CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
PRAI US 1999-133438
                     P 19990511
     This invention relates to cDNAs encoding corn and wheat gene
     RAD51 recombinases. The invention also relates to the
     construction of a chimeric gene encoding all or a substantial portion of
     the recombination protein, in sense or antisense orientation, wherein
     expression of the chimeric gene results in prodn. of altered levels of the
     recombination protein in a transformed host cell. Also disclosed is use
     of the chimeric gene for pos. selection of transformed plant
L_2
     ANSWER 2 OF 49 CAPLUS COPYRIGHT 2001 ACS
ΑN
     2001:397024 CAPLUS
DN
     135:1212
TΙ
     Homologous recombination and molecular evolution of recombination protein
     homologs in plants
IN
     Lassner, Michael; Delcardayre, Steven
PΑ
     Maxygen, Inc., USA
so
     PCT Int. Appl., 57 pp.
     CODEN: PIXXD2
DТ
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                      KIND DATE
                                           APPLICATION NO. DATE
                            _____
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             CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR,
             HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT,
             LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU,
             SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN,
             YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
         RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,
             DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF,
             BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
PRAI US 1999-167450
                      Р
                            19991123
     Methods for evolving recombinase protein homologs and RecA/VirE2 fusion
     proteins which complement VirE2 deficient Agrobacterium are provided.
     use of recombinase protein homologs and RecA/VirE2 fusion proteins in the
     context of Agrobacterium-mediated transformation are provided. Methods
     for producing transgenic organisms by homologous recombination using
     evolved recombinase proteins and Agrobacterium strains which express
     recombinase protein homologs or RecA/VirE2 fusion proteins are provided.
     Transgenic cells and organisms which have integrated an exogenous DNA
     sequence into a predetd. site in their genome are provided.
L2
     ANSWER 5 OF 49 AGRICOLA
                                                         DUPLICATE 1
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ΑN

DN

2001:31834 AGRICOLA

IND22435110

TΤ RAD50 function is essential for telomere maintenance in Arabidopsis. AU Gallego, M.E.; White, C.I. DNAL (500 N21P) ΑV SO Proceedings of the National Academy of Sciences of the United States of America, Feb 13, 2001. Vol. 98, No. 4. p. 1711-1716 Publisher: Washington, D.C.: National Academy of Sciences, CODEN: PNASA6; ISSN: 0027-8424 NTE Includes references District of Columbia; United States CY Article; Conference DT U.S. Imprints not USDA, Experiment or Extension FS T.A English => s (DNA repair) and transform? and (plant or maize or wheat or tobacco or arabidopsis or rice or sunflower) 132 (DNA REPAIR) AND TRANSFORM? AND (PLANT OR MAIZE OR WHEAT OR TOBACCO OR ARABIDOPSIS OR RICE OR SUNFLOWER) => duplicate remove 13 DUPLICATE PREFERENCE IS 'AGRICOLA, BIOSIS, CAPLUS, CABA' KEEP DUPLICATES FROM MORE THAN ONE FILE? Y/(N):n PROCESSING COMPLETED FOR L3 84 DUPLICATE REMOVE L3 (48 DUPLICATES REMOVED) L4=> d ti 1-10 ANSWER 1 OF 84 CAPLUS COPYRIGHT 2001 ACS L4TΤ Methods and uses thereof for generating hypermutable yeast for mutagenesis ANSWER 2 OF 84 CAPLUS COPYRIGHT 2001 ACS T.4 TI Blockade of mismatch  ${\bf DNA}\ {\bf repair}$  in plants and the generation of hypermutable strains for plant breeding L4ANSWER 3 OF 84 CAPLUS COPYRIGHT 2001 ACS Methods of determining individual hypersensitivity to a pharmaceutical TТ agent from gene expression profile ANSWER 4 OF 84 CAPLUS COPYRIGHT 2001 ACS L4ΤT Cell-free assay for plant gene targeting and conversion ANSWER 5 OF 84 CAPLUS COPYRIGHT 2001 ACS TΤ Protein and cDNA sequences of maize retinoblastoma-associatedlike proteins (MSI-like) and uses thereof in enhancing plant disease resistance ANSWER 6 OF 84 CAPLUS COPYRIGHT 2001 ACS L4Uncoupling of 3'-phosphatase and 5'-kinase functions in budding yeast. TΙ Characterization of Saccharomyces cerevisiae DNA 3'-phosphatase (TPP1) ANSWER 7 OF 84 CAPLUS COPYRIGHT 2001 ACS L4Mechanisms of N-acetylcysteine in the prevention of DNA damage and cancer, TΤ with special reference to smoking-related end-points ANSWER 8 OF 84 BIOSIS COPYRIGHT 2001 BIOSIS L4 $\textbf{Arabidopsis} \ \, \textbf{UVH3} \ \, \textbf{gene is a homolog of the Saccharomyces}$ TΤ cerevisiae RAD2 and human XPG DNA repair genes. 1.4 ANSWER 9 OF 84 CAPLUS COPYRIGHT 2001 ACS TΤ Transcriptional activator nucleic acids and polypeptides from plants ANSWER 10 OF 84 CAPLUS COPYRIGHT 2001 ACS TI sequence of  ${\bf Maize}$  replication protein a large and middle subunits with applications for modulation of cell cycle in both dicots and monocots => d ti 11-20

DUPLICATE 2

ANSWER 11 OF 84 CABA COPYRIGHT 2001 CABI

ANSWER 12 OF 84 AGRICOLA

transformed by Agrobacterium.

UV-damage-mediated induction of homologous recombination in **Arabidopsis** is dependent on photosynthetically active radiation.

RecA stimulates sister chromatid exchange and the fidelity of double-strand break repair, but not gene targeting, in plants

TΤ

L4 TI

- L4 ANSWER 13 OF 84 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 3
- TI Competence of **Arabidopsis** thaliana genotypes and mutants for Agrobacterium tumefaciens-mediated gene transfer: Role of phytohormones.
- L4 ANSWER 14 OF 84 CABA COPYRIGHT 2001 CABI
- TI The C termini of **Arabidopsis** cryptochromes mediate a constitutive light response.
- L4 ANSWER 15 OF 84 BIOSIS COPYRIGHT 2001 BIOSIS
- ${\tt TI}$  Factors affecting  ${\tt transformation}$  of Agrobacterium tumefaciens and their application on cereals.
- L4 ANSWER 16 OF 84 AGRICOLA
- TI The **Arabidopsis** UVH1 gene is a homolog of the yeast repair endonuclease RAD1.
- L4 ANSWER 17 OF 84 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 4
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  Arabidopsis UVH1 DNA repair gene is a homolog
  of Saccharomyces cerevisiae Rad1.
- L4 ANSWER 18 OF 84 CAPLUS COPYRIGHT 2001 ACS
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- L4 ANSWER 19 OF 84 CAPLUS COPYRIGHT 2001 ACS
- TI Hijacking oocyte DNA repair machinery in transgenesis?
- L4 ANSWER 20 OF 84 CAPLUS COPYRIGHT 2001 ACS DUPLICATE 5
- TI Integration of T-DNA into plant genomes: prototype and realities
- => d bib abs 12
- L4 ANSWER 12 OF 84 AGRICOLA

DUPLICATÈ 2

- AN 2000:66747 AGRICOLA
- DN IND22063096
- TI RecA stimulates sister chromatid exchange and the fidelity of double-strand break repair, but not gene targeting, in plants transformed by Agrobacterium.
- AU Reiss, B.; Schubert, I.; Kopchen, K.; Wendeler, E.; Schell, J.; Puchta, H.
- AV DNAL (500 N21P)
- SO Proceedings of the National Academy of Sciences of the United States of America, Mar 28, 2000. Vol. 97, No. 7. p. 3358-3363
  Publisher: Washington, D.C.: National Academy of Sciences, CODEN: PNASA6; ISSN: 0027-8424
- NTE Includes references
- CY District of Columbia; United States
- DT Article; Conference
- FS U.S. Imprints not USDA, Experiment or Extension
- LA English
- Expression of the bacterial RecA protein in plants stimulates homologous recombination in tobacco. Here we show that RecA plays a direct role in DNA strand exchange in vivo. The number of sister chromatid exchanges (SCEs) was increased 2.4-fold over wild type in transgenic tobacco plants expressing a nuclear-targeted RecA (nt-RecA) protein and could not be increased further by DNA damage, which caused a doubling of the baseline SCE frequency in wild-type plants. Although gene targeting requires homologous recombination, the number of targeted gene replacements was not increased markedly by the presence of nt-RecA by using Agrobacterium-mediated transformation. However, the number of double-strand breaks that were repaired at both sides by homologous recombination was increased 3.3-fold. Stimulation of SCE and fidelity of double-strand break repair by nt-RecA, but not by gene targeting, suggests that the stimulatory activity of RecA is linked to active DNA synthesis. Therefore, nascent replication-associated single strands may be a prerequisite for RecA action in plant cells.
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- L4 ANSWER 21 OF 84 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 6
- TI Prevention of cancer and other chronic diseases worldwide based on sound mechanisms.
- L4 ANSWER 22 OF 84 CAPLUS COPYRIGHT 2001 ACS
- TI Genetic instability and oral cancer
- L4 ANSWER 23 OF 84 CAPLUS COPYRIGHT 2001 ACS

- TI Solid phase selection of differentially expressed genes by competitive hybridization with reference DNA cloned on microparticles
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- L4 ANSWER 31 OF 84 CAPLUS COPYRIGHT 2001 ACS
- TI Frequent allelic loss at chromosome 3p distinct from genetic alterations of the 8-oxoguanine DNA glycosylase 1 gene in head and neck cancer  $\frac{1}{2}$
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- TI Generation of DNA double-strand breaks and inhibition of somatic embryogenesis by tungsten microparticles in **wheat**.
- L4 ANSWER 34 OF 84 CABA COPYRIGHT 2001 CABI
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- $\ensuremath{\mathsf{TI}}$  . In situ modification of  $\ensuremath{\mathsf{plant}}$  genes for improved herbicide resistance
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- AN 1997:205241 CAPLUS

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     gene disruption to allow exogenous diverse DNA introduction with
     homologous recombination
     Te Riele, Henricus Petrus Josep; De Wind, Niels; Dekker-Vlaar, Helena
ΤN
     Maria Joha
PA
     Setratech, Fr.; Te Riele, Henricus, Petrus, Joseph; De Wind, Niels;
     Dekker-Vlaar, Helena, Maria, Johanna
     PCT Int. Appl., 47 pp.
SO
     CODEN: PIXXD2
DT
     Patent
     English
I.A
FAN.CNT 1
     PATENT NO.
                       KIND DATE
                                             APPLICATION NO. DATE
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                      A1 19970213
     WO 9705268
PΙ
                                             WO 1995-EP2980
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         W: AU, BR, CA, CN, JP, KR, MX, NO, NZ, RU, SE, SG, US
RW: AT, BE, CH, DE, DK, ES, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE,
            BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG
                    A1 19970226
A1 19980520
                                                             19950726
19950726
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     EP 842289
                                             EP 1995-944705
         R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE
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CS
     Weizmann Institute of Science, Rehovot, Israel.
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ΑN
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ΑIJ
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     ANSWER 44 OF 84 BIOSIS COPYRIGHT 2001 BIOSIS DUPLICATE 14
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     expressing bacteriophage T4 DenV and Saccharomyces cerevisiae Apn1
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     (1) Dep. Mol. Biol. and Genet., Univ. Guelph, Guelph, ON N1G 2W1 Canada
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SO
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     English
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     repair and mitotic recombination
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     Hearst Microbiology Research Center, Department of Microbiology, Box 62, Cornell University Medical College, 1300 York Ave., New York, NY 10021,
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     R.L.; Cashmore, A.R.
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     Maryland; United States
CY
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     U.S. Imprints not USDA, Experiment or Extension
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T.4
     ANSWER 47 OF 84 CAPLUS COPYRIGHT 2001 ACS
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     Cancer Research Center of Hawaii, University of Hawaii, Honolulu, HI,
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     genomic double-strand breaks by homologous recombination.
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     Puchta, H.; Dujon, B.; Hohn, B.
CS
     Friedrich Miescher-Institut, Basel, Switzerland.
ΑV
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     Proceedings of the National Academy of Sciences of the United States of
     America, May 14, 1996. Vol. 93, No. 10. p. 5055-5060
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     English
     ANSWER 50 OF 84 AGRICOLA
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     Reiss, B.; Klemm, M.; Kosak, H.; Schell, J.
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DT
     Article; Conference
     U.S. Imprints not USDA, Experiment or Extension
LΑ
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English

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- => file agricola biosis caplus caba
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- L1 2770253 RAD?
- => s rad51?
- L2 1649 RAD51?
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- => duplicate remove 13
- L4 120 DUPLICATE REMOVE L3 (29 DUPLICATES REMOVED)
- => d ti 1-50
- L4 ANSWER 1 OF 120 CAPLUS COPYRIGHT 2003 ACS
- TI Replicative in vivo gene targeting
- L4 ANSWER 2 OF 120 CAPLUS COPYRIGHT 2003 ACS
- $\ensuremath{\mathsf{TI}}$   $\ensuremath{\mathsf{Modulation}}$  of meiotic homologous recombination between non-sister chromatids in meiosis
- L4 ANSWER 3 OF 120 CAPLUS COPYRIGHT 2003 ACS
- TI Methods for enhancing targeted gene alteration in cells having altered activity of DNA repair proteins using chimeric RNA-DNA double-stranded hairpin oligonucleotides
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- L4 ANSWER 5 OF 120 CAPLUS COPYRIGHT 2003 ACS
- TI Cis-acting DNA damage-responsive elements of **Arabidopsis** thaliana **RAD51** gene promoter
- L4 ANSWER 6 OF 120 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.
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- DUPLICATE 1
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- L4 ANSWER 8 OF 120 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.DUPLICATE 2
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- TI BRCA2 homolog required for proficiency in DNA repair, recombination, and genome stability in Ustilago maydis.
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- TI Telomere maintenance without telomerase
- L4 ANSWER 12 OF 120 CAPLUS COPYRIGHT 2003 ACS
- ${
  m TI}$  Homologous recombination induced by replication inhibition, is stimulated by expression of mutant p53
- L4 ANSWER 13 OF 120 CAPLUS COPYRIGHT 2003 ACS
- TI Two rice DMC1 genes are differentially expressed during meiosis and during haploid and diploid mitosis
- L4 ANSWER 14 OF 120 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.

- TI The role of DNA repair in nitrogen mustard drug resistance.
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- L4 ANSWER 18 OF 120 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.
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- TI Antisense-augmented radiotherapy of malignant gliomas
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- TI Assembly of RecA-like recombinases: distinct roles for mediator proteins in mitosis and meiosis.
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- AN 2002:614682 BIOSIS
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- AU Masson, Jean-Yves (1); West, Stephen C. (1)
- CS (1) Imperial Cancer Research Fund, Clare Hall Laboratories, South Mimms, Herts, EN6 3LD: s.west@icrf.icnet.uk UK
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- DT General Review
- LA English
- AB A double-strand break in genomic DNA that remains unrepaired can be lethal for a cell. Indeed, the integrity of the genome is paramount for survival. It is therefore surprising that some cells deliberately introduce double-strand breaks at certain times during their life cycle. Why might they do this? What are the benefits? How are these breaks repaired? The

answers to these questions lie in understanding the basis of meiotic recombination, the process that leads to genetic variation. This **review** summarizes the key roles played by the two recombinases, Dmcl and **Rad51**, in the faithful repair of DNA breaks.

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ANSWER 44 OF 120 CAPLUS COPYRIGHT 2003 ACS
1.4
AN
     2000:814630 CAPLUS
     133:360459
DN
     Gene RAD51 recombinase and cDNA of corn and wheat and
TΙ
     use of RAD51 in positive selection of recombinant plant
IN
     Famodu, Omolayo O.; Morgante, Michele
     E.I. du Pont de Nemours and Company, USA
PΑ
SO
     PCT Int. Appl., 46 pp.
     CODEN: PIXXD2
DТ
     Patent
LA
     English
FAN.CNT 1
     PATENT NO.
                       KIND DATE
                                             APPLICATION NO. DATE
                             -----
PΙ
     WO 2000068390
                       A2
                             20001116
                                             WO 2000-US12587 20000509
     WO 2000068390
                       A3
                             20010503
         W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU,
             CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL,
             IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD,
             MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK,
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         RW: GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF,
              CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
     US 2002152497
                        A1
                            20021017
                                             US 2002-78929
                                                               20020219
PRAI US 1999-133438P
                             19990511
                        Ρ
     US 1999-133038P
                             19990507
     US 1999-133042P
                        Ρ
                             19990507
     US 1999-133427P
                        Р
                             19990511
     US 1999-133428P
                        Ρ
                             19990511
     US 1999-133436P
                        Р
                             19990511
     US 1999-133437P
                        Ρ
                             19990511
                       P
A3
     US 1999-137667P
                             19990604
     US 2000-566394
                             20000505
AB
     This invention relates to cDNAs encoding corn and wheat gene
     RAD51 recombinases. The invention also relates to the
     construction of a chimeric gene encoding all or a substantial portion of
     the recombination protein, in sense or antisense orientation, wherein
     expression of the chimeric gene results in prodn. of altered levels of the
     recombination protein in a transformed host cell. Also disclosed is use
     of the chimeric gene for pos. selection of transformed plant
     cells.
     ANSWER 22 OF 120 CAPLUS COPYRIGHT 2003 ACS
     2001:397024 CAPLUS
ΑN
DN
     135:1212
ΤT
     Homologous recombination and molecular evolution of recombination protein
     homologs in plants
ΙN
     Lassner, Michael; Delcardayre, Steven
PΑ
     Maxygen, Inc., USA
SO
     PCT Int. Appl., 57 pp.
     CODEN: PIXXD2
DT
     Patent
T.A
     English
FAN.CNT 1
     PATENT NO.
                       KIND DATE
                                             APPLICATION NO. DATE
     WO 2001038504
                        A2
                              20010531
                                             WO 2000-US32289 20001122
PΙ
     WO 2001038504
                        АЗ
                             20020124
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              HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT,
              LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU,
              SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN,
         YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW; AT, BE, CH, CY,
              DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF,
              BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
167450P P 19991123
PRAI US 1999-167450P
     Methods for evolving recombinase protein homologs and RecA/VirE2 fusion
     proteins which complement VirE2 deficient Agrobacterium are provided. The
     use of recombinase protein homologs and RecA/VirE2 fusion proteins in the
     context of Agrobacterium-mediated transformation are provided. Methods
     for producing transgenic organisms by homologous recombination using
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evolved recombinase proteins and Agrobacterium strains which express

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recombinase protein homologs or RecA/VirE2 fusion proteins are provided. Transgenic cells and organisms which have integrated an exogenous DNA sequence into a predetd. site in their genome are provided.

- L4 ANSWER 13 OF 120 CAPLUS COPYRIGHT 2003 ACS
- AN 2002:243417 CAPLUS
- DN 137:258167
- TI Two rice DMC1 genes are differentially expressed during meiosis and during haploid and diploid mitosis
- AU Kathiresan, Arumugam; Khush, Gurdev S.; Bennett, John
- CS Plant Breeding, Genetics and Biochemistry Division, International Rice Research Institute, Metro Manila, Philippines
- SO Sexual Plant Reproduction (2002), 14(5), 257-267 CODEN: SPLRE7; ISSN: 0934-0882
- PB Springer-Verlag
- DT Journal
- LA English
- We have cloned two **rice** homologues of yeast DMC1, a meiosis-specific gene required for recombination between homologous chromosomes. We show that **rice** DMC1A and DMC1B were produced by a gene duplication event that occurred after **rice** sepd. from the common ancestor of the cereals. The predicted proteins contain 344 amino acids, of which all but 7 are conserved between the two homologues. Between bases -1 and -245, the two promoters share six invariant blocks of sequence of 10-28 bp, interspersed in variable sequences. Both DMC1A and DMC1B are expressed in pollen mother cells coincident with meiosis, and in diploid non-meiotic tissues such as calli and root tips. DMC1B is also expressed in haploid male gametophytes during pollen maturation and in diploid zygotic embryos and endosperm after pollination. These data suggest that DMC1B, either alone or in combination with DMC1A, contributes to recombination during meiosis and during haploid and diploid mitosis.
- RE.CNT 40 THERE ARE 40 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT
- L4 ANSWER 15 OF 120 BIOSIS COPYRIGHT 2003 BIOLOGICAL ABSTRACTS INC.
  DUPLICATE 4
- AN 2002:451682 BIOSIS
- DN PREV200200451682
- TI Molecular cloning and characterization of RAD51-like genes from Arabidopsis thaliana.
- AU Osakabe, Keishi; Yoshioka, Toji; Ichikawa, Hiroaki; Toki, Seiichi (1)
- CS (1) Department of Plant Biotechnology, National Institute of Agrobiological Sciences, 2-1-2 Kannondai, Tsukuba, Ibaraki, 305-8602: stoki@affrc.go.jp Japan
- SO Plant Molecular Biology, (September, 2002) Vol. 50, No. 1, pp. 71-81. http://www.kluweronline.com/issn/0167-4412. print. ISSN: 0167-4412.
- DT Article
- LA English
- AB Homologous recombination is an essential process for the maintenance and variability of the genome. In eukaryotes, the Rad52 epistasis group proteins serve the main role for meiotic recombination and/or homologous recombinational repair. Rad51-like proteins, such as Rad55 and Rad57 in yeast, play a role in assembly or stabilization of multimeric Rad51 that promotes homologous pairing and strand exchange reactions. We cloned two RAD51-like genes named AtXRCC3 and AtRAD51C from Arabidopsis thaliana. Both AtXRCC3 and AtRAD51C expressed two alternatively spliced transcripts, and AtRAD51C produced two different sizes of isoforms, a long (AtRAD51Calpha) and a short one (AtRAD51Cbeta). The predicted protein sequences of these genes showed characteristic features of the RecA/Rad51 family; especially the amino acids around the ATP-binding motifs were well conserved. The transcripts of AtXRCC3 and AtRAD51C were detected in various tissues, with the highest level of expression in flower buds. Expression of both genes was induced by gamma-ray irradiation. The results of yeast two-hybrid assays suggested that Arabidopsis Rad51 family proteins form a complex, which could participate in meiotic recombination and/or homologous recombinational repair.

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